## Output tables for the test of Multiple comparisons.

December 17, 2022

## 1 Average rankings of Friedman test

Average ranks obtained by applying the Friedman procedure

Ranking	6.1034	3.9655	4.7586	3.5172	4.8621	3.2759	1.5172
Algorithm	MEABC	ABCADE	ABCNG	SHADE	MAPSO	TAPSO	MEEABC

Table 1: Average Rankings of the algorithms

Friedman statistic considering reduction performance (distributed according to chi-square with 6 degrees of freedom: 78.70936. P-value computed by Friedman Test: 4.016320609423474E-11.

## 2 Post hoc comparisons

Results achieved on post hoc comparisons for  $\alpha=0.05,\,\alpha=0.10$  and adjusted p-values.

## 2.1 P-values for $\alpha = 0.05$

i	algorithms	$z = (R_0 - R_i)/SE$	d	Holm
21	MEABC vs. MEEABC	8.084148	0	0.002381
20	MAPSO vs. MEEABC	5.895958	0	0.0025
19	ABCNG vs. MEEABC	5.713608	0	0.002632
18	MEABC vs. TAPSO	4.984212	0.000001	0.002778
17	MEABC vs. SHADE	4.55873	0.000005	0.002941
16	ABCADE vs. MEEABC	4.315598	0.000016	0.003125
15	MEABC vs. ABCADE	3.76855	0.000164	0.003333
14	SHADE vs. MEEABC	3.525418	0.000423	0.003571
13	TAPSO vs. MEEABC	3.099936	0.001936	0.003846
12	MAPSO vs. TAPSO	2.796021	0.005174	0.004167
11	ABCNG vs. TAPSO	2.613672	0.008958	0.004545
10	MEABC vs. ABCNG	2.37054	0.017762	0.005
6	SHADE vs. MAPSO	2.37054	0.017762	0.005556
œ	MEABC vs. MAPSO	2.18819	0.028656	0.00625
_	ABCNG vs. SHADE	2.18819	0.028656	0.007143
9	ABCADE vs. MAPSO	1.58036	0.114025	0.008333
20	ABCADE vs. ABCNG	1.398011	0.16211	0.01
4	ABCADE vs. TAPSO	1.215661	0.224114	0.0125
က	ABCADE vs. SHADE	0.79018	0.429423	0.016667
2	SHADE vs. TAPSO	0.425481	0.670486	0.025
1	ABCNG vs. MAPSO	0.182349	0.855309	0.05

Table 2: P-values Table for  $\alpha = 0.05$ 

Holm's procedure rejects those hypotheses that have an unadjusted p-value  $\leq 0.004167$ .

$z = \frac{(R_0 - R_i)/SE}{8.084148}$
5.895958
5.713608
4.984212
4.55873
4.315598
3.76855
3.525418
3.099936
2.796021
2.613672
2.37054
2.37054
2.18819
2.18819
1.58036
1.39801
1.215661
0.79018
0.425481
0.182349

Table 3: P-values Table for  $\alpha = 0.10$ 

Holm's procedure rejects those hypotheses that have an unadjusted p-value  $\leq 0.01$ .

$p_{Holm}$	0	0	0	0.000011	0.000087	0.000255	0.002463	0.005919	0.025163	0.062083	0.098533	0.177621	0.177621	0.229246	0.229246	0.684147	0.810549	0.896456	1.288268	1.340971	1.340971
unadjusted $p$	0	0	0	0.000001	0.000005	0.000016	0.000164	0.000423	0.001936	0.005174	0.008958	0.017762	0.017762	0.028656	0.028656	0.114025	0.16211	0.224114	0.429423	0.670486	0.855309
hypothesis	MEABC vs .MEEABC	MAPSO vs .MEEABC	ABCNG vs .MEEABC	MEABC vs .TAPSO	MEABC vs .SHADE	ABCADE vs .MEEABC	MEABC vs .ABCADE	SHADE vs .MEEABC	TAPSO vs .MEEABC	MAPSO vs .TAPSO	ABCNG vs .TAPSO	MEABC vs .ABCNG	SHADE vs .MAPSO	MEABC vs .MAPSO	ABCNG vs .SHADE	ABCADE vs .MAPSO	ABCADE vs .ABCNG	ABCADE vs .TAPSO	ABCADE vs .SHADE	SHADE vs .TAPSO	ABCNG vs .MAPSO
	1	2	33	4	ಬ	9	-1	<sub>∞</sub>	6	10	11	12	13	14	15	16	17	18	19	20	21

Table 4: Adjusted p-values